

SEQUENCE LISTING

<110> UAB Research Foundation

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CHAUDHURI, Tandra R.
WU, Hongju

<120> METHODS AND COMPOSITIONS FOR IN VIVO
INFLAMMATION MONITORING

<130> 21085.0050P1

<140> Unassigned

<141> 2004-09-23

<150> 60/505,543

<151> 2003-09-23

<160> 29

<170> FastSEQ for Windows Version 4.0

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<213> Artificial Sequence

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<223> Description of Artificial Sequence; note =
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aagcacatcg tagtcatgct catgcagata aaggcaggta agctccggaa ccaccacaga 35160
aaaagacacc atttttctct caaacatgtc tgcgggtttc tgcataaaca caaaataaaaa 35220
taacaaaaaa acattttaaac attagaagcc tgtcttacaa caggaaaaac aacccttata 35280
agcataagac ggactacggc catgccggcg tgaccgtaaa aaaactgggc accgtgatta 35340
aaaagcacca ccgacagctc ctccggtcatg tccggagtca taatgtaaga ctccgtaaac 35400
acatcagggtt gattcatcgg tcagtgtctaa aaagcgaccg aaatagcccg ggggaatata 35460
taccgcgagg cgtagagaca acattacagc ccccatagga ggtataacaa aattaatagg 35520
agagaaaaac acataaacac ctgaaaaacc ctctgccta ggcaaaatag caccctcccg 35580
ctccagaaca acatacagcg cttcacagcg gcagcctaac agtcagcctt accagtaaaa 35640
aagaaaacct attaaaaaaa caccactcga cacggcacca gctcaatcag tcacagtgtg 35700
aaaaagggcc aagtgcagag cgagtatata taggactaaa aaatgacgta acggttaaaag 35760
tccacaaaaa acaccagaa aaccgcacgc gaacctacgc ccagaaacga aagccaaaaa 35820
accacaact tcctcaaatc gtcacttcgg ttttcccacg ttacgtaact tcccatttta 35880
agaaaactac aattcccaac acatacaagt tactccggcc taaaacctac gtcacccgcc 35940
ccgttcccac gcccgcgcc acgtcacaaa ctccaccccc tcattatcat attggcttca 36000
atccaaaata aggtatatta ttgatgatg 36029

```

<210> 6

<211> 720

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<221> misc_feature

<222> 693,709

<223> n = g, a, c or t(u)

<400> 6

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ttcaactagg tgtcctcgga tcccacgaag tgaaaattaa acacttttct ccgtatcacg 60
aagtgaaaat taaacacttt tctccgtatg gatcccatca ccatcaccat cacctagggt 120
cacctaaata tgccgataaa acatttcaac ctgaacctca aataggagaa tctcagtggg 180
acgaaacaga aattaatcat gcagctggga gagtcctaaa aaagactacc ccaatgaaac 240
catgttacgg ttcatatgca aaacccacaa atgaaaatgg agggcaaggc attcttgtaa 300
agcaacaaaa tggaaagcta gaaagtcaag tggaaatgca atttttctca actactgagg 360
cagccgcagg caatgggtgat aacttgactc ctaaagtggg attgtacagt gaagatgtag 420
atatagaaac ccagacact catatttctt acatgcccac tattaaggaa ggtaactcac 480
gagaactaat gggocaacaa tctatgccca acaggcctaa ttacattgct tttagggaca 540
atthttattg tctaattgat tacaacagca cgggtaatat ggggtgttctg gcggggccaag 600
catcgcagtt gaatgctgtt gtagatttgc aagacagaaa cacagagctt tcataccagc 660
ttttgcttga ttccattggg gatagaacca ggntactttt ctatgtggna tcaggctggg 720

```

<210> 7

<211> 719

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<221> misc_feature

<222> 2,3,4,10,12,16,17,42,715,719

<223> n = g, a, c or t(u)

<400> 7

```

cnnnggaggn cnttcnnata ggtgtcgaag gtcaaacacc tnaatatgcc gataaaacat      60
ttcaacctga acctcaaata ggagaatctc agtggtacga aacagaaatt aatcatgcag      120
ctgggagagt cctaaaaaag actaccccaa tgaaaccatg ttacggttca tatgcaaaac      180
ccacaaatga aaatggaggg caaggcattc ttgtaaagca acaaaatgga aagctagaaa      240
gtcaagtgga aatgcaatth ttctcaacta ctctcggatc ccacgaagtg aaaattaaac      300
acttttctcc gtatcacgaa gtgaaaatta aacacttttc tccgtatgga tcccatcacc      360
atcaccatca cctaggttca ttgactccta aagtgggtatt gtacagtga gatgtagata      420
tagaaacccc agacactcat atttcttaca tgcccactat taaggaagggt aactcacgag      480
aactaatggg ccaacaatct atgcccaaca ggcctaatta cattgctttt agggacaatt      540
ttattgggtc aatgtattac aacagcacgg gtaatatggg tgttctggcg ggccaagcat      600
cgcagttgaa tgctgttgta gatttgcaag acagaaacac agagctttca taccagcttt      660
tgcttgattc cattgggtgat agaaccagggt acttttctat gtggaatcag gctgntgan      719

```

<210> 8

<211> 108

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 8

```

ctcggatccc acgaagtga aattaaacac ttttctccgt atcacgaagt gaaaattaaa      60
cacttttctc cgtatggatc ccatcaccat caccatcacc taggttca      108

```

<210> 9

<211> 36

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 9

```

Leu Gly Ser His Glu Val Lys Ile Lys His Phe Ser Pro Tyr His Glu
 1             5             10            15
Val Lys Ile Lys His Phe Ser Pro Tyr Gly Ser His His His His
      20             25             30

```

```

His Leu Gly Ser
      35

```

<210> 10

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 10

```

His Glu Val Lys Ile Lys His Phe Ser Pro Tyr
 1             5             10

```

<210> 11

<211> 5

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 11

Gly Gly Gly Gly Ser
1 5

<210> 12

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 12

Leu Gly Ser His His His His His His Leu Gly Ser
1 5 10

<210> 13

<211> 3

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 13

Lys Gly Ser
1

<210> 14

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 14

cctacgcacg acgtgaccac ag

22

<210> 15

<211> 62

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 15

tgaacctagg tgatggtgat ggtgatggga tccgaggaca cctatttgaa taccctcctt
tg

60

62

<210> 16
 <211> 61
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 16
 ctcgatccc atcaccatca ccatcaccta gggtcaccta aatatgcga taaaacattt 60
 c 61

 <210> 17
 <211> 23
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 17
 ctagggagct ctgcagaacc atg 23

 <210> 18
 <211> 60
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 18
 tgaacctagg tgatggtgat ggtgatggga tccgagttcg taccactgag attctcctat 60

 <210> 19
 <211> 60
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 19
 ctcgatccc atcaccatca ccatcaccta gggtcaactg aaattaatca tgcagctggg 60

 <210> 20
 <211> 61
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 20
 tgaacctagg tgatggtgat ggtgatggga tccgagagta gttgagaaaa attgcatttc 60
 c 61

<210> 21
 <211> 60
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 21
 ctcggatccc atcaccatca ccatcaccta ggttcattga ctcctaaagt ggtattgtac 60

 <210> 22
 <211> 61
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 22
 tgaacctagg tgatggtgat ggtgatggga tccgagagtg ggcattgtaag aaatatgagt 60
 g 61

 <210> 23
 <211> 58
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 23
 ctcggatccc atcaccatca ccatcaccta ggttcaaact cacgagaact aatggggc 58

 <210> 24
 <211> 60
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 24
 tgaacctagg tgatggtgat ggtgatggga tccgagaggt ttaccttg taagagtctc 60

 <210> 25
 <211> 62
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Description of Artificial Sequence; note =
 synthetic construct

 <400> 25
 ctcggatccc atcaccatca ccatcaccta ggttcattggg aaaaagatgc tacagaattt 60

tc

62

<210> 26

<211> 60

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 26

tgaacctagg tgatggtgat ggtgatggga tccgagtgga aagcagtaat ttggaagttc 60

<210> 27

<211> 63

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 27

ctcggatccc atcaccatca ccatcaccta ggttcaaata attttgccat ggaaatcaat 60
cta 63

<210> 28

<211> 720

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<221> misc_feature

<222> 1-14, 123-720

<223> n = g, a, c or t(u)

<400> 28

```

nnnnnnnnnn nnnnctcgga tcccacgaag tgaaaattaa acacttttct ccgtatcacg      60
aagtgtaaaat taaacacttt tctccgtatg gatcccatca ccatcaccat cacctagggtt    120
cannnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    180
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    240
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    300
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    360
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    420
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    480
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    540
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    600
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    660
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn    720

```

<210> 29

<211> 720

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =

synthetic construct

<221> misc_feature

<222> 1-282, 390-720

<223> n = g, a, c or t(u)

<400> 29

nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	60
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	120
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	180
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	240
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnctcggatc	ccacgaagtg	aaaattaaac	300
acttttctcc	gtatcacgaa	gtgaaaatta	aacacttttc	tccgtatgga	tcccatcacc	360
atcaccatca	cctaggttca	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	420
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	480
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	540
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	600
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	660
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	720